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SEQUENCE LISTING

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<120> HIGH-THROUGHPUT SCREENING OF EXPRESSED DNA LIBRARIES IN
FILAMENTOUS FUNGI

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<141> 2001-04-13

<150> PCT/US00/10199

<151> 2000-04-13

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<170> PatentIn Ver. 2.1

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 35 40 45
 Ala His Tyr Ser Gln Ser Ile Leu Val His Thr Ala Phe Gly Cys Gly
 50 55 60
 Val Leu Thr Ser Ser Thr Arg Met Ser Pro Thr Phe Leu Ser Gln Ser
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 Ile Ile Ala Ser Lys Phe Pro Arg Asn Phe Pro Leu Gln Pro Arg Val
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 Thr Arg Pro Pro Ser Trp Ser Leu Ser Ser Ala Asn Val Leu Thr Phe
 115 120 125
 Gly Thr Phe Thr Leu Lys Ser Gly Arg Arg Ala Ser Pro Leu Gln His
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 145 150 155 160
 Thr Ser Ser Thr Pro Ala Ser Ser Thr Pro Pro Leu Ser Ser Pro Pro
 165 170 175
 Ser Pro Pro Trp Pro Thr Pro Ser Ser Pro Ser Ser Leu Arg Thr Leu
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 Pro Ser Pro Ser Pro Thr Ser Cys Phe Gly Lys Thr Pro Ser Phe Pro
 195 200 205
 Asn Thr Pro Leu Pro Leu Asn Asn Pro Ile Thr Asn Lys Asn Pro Leu
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 Tyr Asn Arg Lys Glu Ala Lys Asp His Gly Glu Gly Gly Asn Ile Val
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Gly Ala Ala Leu Lys Gly Lys Thr Val Leu Val Ile Asp Asp Val Ile
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 Asn Glu Ala Asp Met Lys Arg Leu Glu Glu Tyr Arg Ala Lys Tyr Gln
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 Met Leu Gly Thr Pro Val Pro Thr Gln Trp Ala Pro Val Ser His Ile
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 His Pro Pro Glu Ala Gln Leu Phe Cys Leu Asn Gln Leu His Pro Asn
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 Val Ala Val Arg His Pro Ser Asn Thr Gly Ile Ile Ala Ile Gly Arg
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 His His Leu Pro Arg Glu Pro Phe His Pro Gln Ala Arg Arg His Ala
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 Ser Gly Lys Lys Pro Pro Leu Ser Pro Ile Pro His Phe His Ser Thr
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 Thr Ala Lys Ala Ala Thr Leu Ser Ala Pro Leu Arg Ala Arg Pro Cys
 260 265 270
 Leu Ser Thr Met Ser Ser Arg Pro Val Pro Pro Cys Val Arg Pro Ser
 275 280 285
 Thr Trp Ser Pro Arg Arg Ala Ala Arg Ser Ser Asp Ser Leu Leu Leu
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 Trp Thr Ala Trp Arg Arg Cys Pro Asp Pro Arg Thr Arg Thr Val Ser
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 370 375 380
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 Pro Lys Arg Pro Leu Gly Phe Ser Ile Ser Pro Glu Arg Cys Ser Ala
 420 425 430
 Arg His Glu Ser Ser Pro Leu Ser Gly His Pro Phe Pro Thr Phe Glu
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 Val Ala Asp Ala Tyr Leu Ala Glu Ala Val Ala Trp Lys Gly Thr Met
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 35 40 45
 Arg Thr Leu Phe Pro Lys His Thr Gly Thr Tyr Cys Ile Arg Leu Val
 50 55 60
 Arg Gly Ala Tyr Leu Ile His Pro Asn Glu Pro Asn Phe Phe Val Ser
 65 70 75 80
 Ile Asn Asn Cys Ile Gln Ile Pro Pro Gln Leu Pro Pro Pro Thr Pro
 85 90 95
 Cys Leu Tyr His Ser Leu His Thr His Thr Ile Thr Met Ala Leu Pro
 100 105 110

Ala Tyr Lys Thr Ala Phe Leu Glu Ser Leu Val Gly Gln Arg Ala Asp
 115 120 125
 Phe Arg His Leu His Pro Glu Val Gly Ser Pro Cys Val Thr Pro Pro
 130 135 140
 Thr Pro Ala Leu Ser Gln Ser Glu Asp Leu Pro Leu Tyr Thr Asp Ser
 145 150 155 160
 Pro Tyr Phe Phe Asn Ala Gly Ile Phe Asn Thr Ala Ser Leu Leu Ser
 165 170 175
 Ala Leu Ser Thr Met Ala His Thr Ile Ile Thr Phe Leu Ala Glu Asn
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 Pro Ser Ile Pro Lys Pro Asp Val Met Leu Arg Val Lys Asn Pro Leu
 195 200 205
 Phe Pro Gln Tyr Pro Thr Ser Thr Gln Gln Pro Ile Asn Asn Gln Lys
 210 215 220
 Pro Pro Lys Gln Pro Arg Ile Gln Arg His Pro Pro Arg Val Arg His
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 Pro Pro Thr Gln Pro His Arg Pro Arg His Leu Gly Gln Arg Val Leu
 245 250 255
 Gln Leu Gln Pro Gln Arg Ser Gln Gly Ser Arg Arg Arg Arg Gln His
 260 265 270
 Cys Arg Arg Arg Ser Glu Gly Gln Asp Arg Ala Cys Asp Arg Arg Cys
 275 280 285
 His His Gly Arg Tyr Arg His Ala Asp Pro Gln Pro Gly Arg Gln Gly
 290 295 300
 Gly Arg Gln Gly Arg Arg Ile His Cys Cys Ser Gly Pro Leu Gly Glu
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 Asp Ala Arg Thr Gln Gly Arg Glu Arg Cys Arg Gly Arg Ala Gln Asn
 325 330 335
 Glu Cys Tyr Gly Ser Asp Pro Gly Val Trp Cys Ala His Asp Glu Tyr
 340 345 350
 Cys Tyr Ser Gly Phe Asp Gln Val Asp Ala Gly Glu Gly Gln Gly Arg
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 Tyr Glu Ala Val Gly Gly Val Gly Val Ser Gly Leu Val Gly Phe Ile
 370 375 380
 Asp Arg Leu Phe Gly Trp Val Glu Val Arg Leu Gly Cys Gly Arg Arg
 385 390 395 400
 Asn Glu Lys Leu Tyr Thr Gly Pro Glu Glu Val Arg Arg Asp Gly Arg
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Glu Met Phe Tyr Val Lys Ile Leu Asn Lys His Leu Lys Lys Asp Pro
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Leu Val Ser Ala Glu Leu Ala Arg Lys Asp Ala Arg His Ala Met Ser
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Leu Ala His Ser Val Gly Thr Arg Phe Pro His Leu Lys Trp Pro Thr
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Leu Ile Trp Leu Arg Leu Trp Pro Gly Lys Ala Leu Trp Arg Ala Ala
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